

A-71864.ST25.txt
SEQUENCE LISTING

<110> Fong, Timothy
Te, Alexis

<120> Cytomodulating Peptides for Treating Interstitial Cystitis

<130> A-71864 (465840-TBD)

<140> To Be Determined

<141> 2005-05-16

<150> PCT/US2003/037043

<151> 2003-11-17

<150> US 60/426,684

<151> 2003-05-15

<150> US 60/426,684

<151> 2002-11-15

<160> 35

<170> PatentIn version 3.3

<210> 1

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<212> PRT

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<223> Synthetic

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<221> MISC_FEATURE

<222> (1)..(1)

<223> The Xaa at position 1 can be any basic amino acid, preferably lysine or arginine

<220>

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<222> (2)..(4)

<223> The Xaa at positions 2 to 4 can be any non-polar aliphatic or aromatic amino acid of from 5 to 6 carbon atoms, preferably any amino acid other than a polar aliphatic amino acid

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<222> (5)..(5)

<223> The Xaa at position 5 can be any basic amino acid, preferably lysine or arginine

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<222> (6)..(8)

<223> The Xaa at positions 6 to 8 can be any non-polar aliphatic or aromatic amino acid of from 5 to 6 carbon atoms, preferably any amino acid other than a polar aliphatic amino acid

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<222> (9)..(9)

<223> The Xaa at position 9 can be glycine, or any basic amino acid, or an aliphatic hydrophobic amino acid of from 5 to 6 carbon atoms

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Tyr
1 5 10

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<223> The Xaa at positions 3 to 4 can be any non-polar aliphatic or aromatic amino acid of from 5 to 6 carbon atoms, preferably any amino acid other than a polar aliphatic amino acid

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<222> (9)..(9)

<223> The Xaa at position 9 can be glycine, or any basic amino acid, or an aliphatic hydrophobic amino acid of from 5 to 6 carbon atoms

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Arg Leu Leu Leu Arg Leu Leu Leu Gly Tyr
1 5 10

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Arg Val Leu Leu Arg Leu Leu Leu Gly Tyr
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Arg Ile Leu Leu Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Val Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Ile Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Leu Arg Val Leu Leu Gly Tyr
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Arg Leu Leu Leu Arg Ile Leu Leu Gly Tyr
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Arg Leu Leu Leu Arg Leu Ile Leu Gly Tyr
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Arg Leu Leu Leu Arg Leu Leu Ile Gly Tyr
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Arg Trp Leu Leu Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Trp Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Leu Arg Leu Trp Leu Gly Tyr
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Arg Leu Leu Leu Arg Leu Leu Trp Gly Tyr
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Arg Tyr Leu Leu Arg Leu Leu Leu Gly Tyr
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Arg Leu Tyr Leu Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Tyr Arg Leu Leu Leu Gly Tyr
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Arg Leu Leu Leu Arg Tyr Leu Leu Gly Tyr
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<223> Synthetic

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Arg Leu Leu Leu Arg Leu Tyr Leu Gly Tyr
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<211> 10

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Arg Leu Leu Leu Arg Leu Leu Tyr Gly Tyr
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<222> (2)..(4)

<223> The Xaa at positions 2 to 4 are norleucine or any D-stereoisomer
amino acid

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<222> (6)..(8)

<223> The Xaa at positions 2 to 4 are norleucine or any D-stereoisomer
amino acid

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Arg Xaa Xaa Xaa Arg Xaa Xaa Xaa Gly Tyr
1 5 10

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<400> 29

Gly Ser Gly Gly Ser
1 5

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Gly Gly Gly Ser
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<222> (1)..(5)
<223> The Xaa at positions 1 to 5 can be any amino acid

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<222> (7)..(9)
<223> The xaa at positions 7 to 9 can be any amino acid, where one of amino acids 7 to 9 can be absent

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<222> (11)..(22)
<223> The Xaa at positions 11 to 22 can be any amino acid, where up to 8 of amino acids 11 to 22 can be absent

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<222> (24)..(26)
<223> The Xaa at positions 24 to 26 can be any amino acid

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<221> MISC_FEATURE
<222> (28)..(32)
<223> The Xaa at positions 28 to 32 can be any amino acid

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Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Xaa
20 25 30

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<222> (7)..(26)
<223> The Xaa at positions 7 to 26 can be any amino acid, where up to 17 amino acids 7 to 26 can be absent

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Phe Gln Cys Glu Glu Cys Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Ile Arg Ser His Thr
20 25 30

Gly

<210> 33
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<222> (2)..(3)
<223> The Xaa at positions 2 to 3 can be any amino acid

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<222> (4)..(24)
<223> The Xaa at positions 4 to 24 can be any amino acid, where up to 16 amino acids 4 to 24 can be absent

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<222> (26)..(29)
<223> The xaa at positions 26 to 29 can be any amino acid

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Cys Xaa Xaa Cys Xaa
1 5 10 15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa Xaa Xaa Cys
20 25 30

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<222> (7)..(26)
<223> The Xaa at positions 7 to 26 can be any amino acid, where up to 16 amino acids 7 to 26 can be absent

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Val Lys Cys Phe Asn Cys Xaa 1
5 10 15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Thr Ala Arg Asn Cys 20
25 30

Arg

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<220>
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<222> (10)..(29)
<223> The Xaa at positions 10 to 29 can be any amino acid, where up to 16 amino acids 10 to 29 can be absent

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Met Asn Pro Asn Cys Ala Arg Cys Gly Xaa Xaa Xaa Xaa Xaa Xaa Xaa 1
5 10 15

Xaa His Lys Ala 20
25 30

Cys Phe